

## SEQUENCE LISTING

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<120> GLUTAMATE RECEPTOR ANTAGONISTS AS NEUROPROTECTIVES

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<150> DE 10337098.6

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<160> 1

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<210> 1

<211> 477

<212> PRT

<213> Desmodus rotundus

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Gly	Ala	Val	Phe	Ser	Leu	Pro	Arg	Gln	Glu	Thr	Tyr	Arg	Gln	Leu	Ala
			20					25					30		

Arg	Gly	Ser	Arg	Ala	Tyr	Gly	Val	Ala	Cys	Lys	Asp	Glu	Ile	Thr	Gln
		35					40					45			

Met	Thr	Tyr	Arg	Arg	Gln	Glu	Ser	Trp	Leu	Arg	Pro	Glu	Val	Arg	Ser
	50					55					60				

Lys	Arg	Val	Glu	His	Cys	Gln	Cys	Asp	Arg	Gly	Gln	Ala	Arg	Cys	His
65					70					75				80	

Thr	Val	Pro	Val	Asn	Ser	Cys	Ser	Glu	Pro	Arg	Cys	Phe	Asn	Gly	Gly
				85					90					95	

Thr	Cys	Trp	Gln	Ala	Val	Tyr	Phe	Ser	Asp	Phe	Val	Cys	Gln	Cys	Pro
			100					105					110		

Ala	Gly	Tyr	Thr	Gly	Lys	Arg	Cys	Glu	Val	Asp	Thr	Arg	Ala	Thr	Cys
							120					125			

Tyr Glu Gly Gln Gly Val Thr Tyr Arg Gly Thr Trp Ser Thr Ala Glu  
 130 135 140  
 Ser Arg Val Glu Cys Ile Asn Trp Asn Ser Ser Leu Leu Thr Arg Arg  
 145 150 155 160  
 Thr Tyr Asn Gly Arg Met Pro Asp Ala Phe Asn Leu Gly Leu Gly Asn  
 165 170 175  
 His Asn Tyr Cys Arg Asn Pro Asn Gly Ala Pro Lys Pro Trp Cys Tyr  
 180 185 190  
 Val Ile Lys Ala Gly Lys Phe Thr Ser Glu Ser Cys Ser Val Pro Val  
 195 200 205  
 Cys Ser Lys Ala Thr Cys Gly Leu Arg Lys Tyr Lys Glu Pro Gln Leu  
 210 215 220  
 His Ser Thr Gly Gly Leu Phe Thr Asp Ile Thr Ser His Pro Trp Gln  
 225 230 235 240  
 Ala Ala Ile Phe Ala Gln Asn Arg Arg Ser Ser Gly Glu Arg Phe Leu  
 245 250 255  
 Cys Gly Gly Ile Leu Ile Ser Ser Cys Trp Val Leu Thr Ala Ala His  
 260 265 270  
 Cys Phe Gln Glu Ser Tyr Leu Pro Asp Gln Leu Lys Val Val Leu Gly  
 275 280 285  
 Arg Thr Tyr Arg Val Lys Pro Gly Glu Glu Glu Gln Thr Phe Lys Val  
 290 295 300  
 Lys Lys Tyr Ile Val His Lys Glu Phe Asp Asp Asp Thr Tyr Asn Asn  
 305 310 315 320  
 Asp Ile Ala Leu Leu Gln Leu Lys Ser Asp Ser Pro Gln Cys Ala Gln  
 325 330 335  
 Glu Ser Asp Ser Val Arg Ala Ile Cys Leu Pro Glu Ala Asn Leu Gln  
 340 345 350  
 Leu Pro Asp Trp Thr Glu Cys Glu Leu Ser Gly Tyr Gly Lys His Lys  
 355 360 365  
 Ser Ser Ser Pro Phe Tyr Ser Glu Gln Leu Lys Glu Gly His Val Arg  
 370 375 380  
 Leu Tyr Pro Ser Ser Arg Cys Ala Pro Lys Phe Leu Phe Asn Lys Thr  
 385 390 395 400  
 Val Thr Asn Asn Met Leu Cys Ala Gly Asp Thr Arg Ser Gly Glu Ile  
 405 410 415  
 Tyr Pro Asn Val His Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu  
 420 425 430

Val Cys Met Asn Asp Asn His Met Thr Leu Leu Gly Ile Ile Ser Trp  
435 440 445

Gly Val Gly Cys Gly Glu Lys Asp Val Pro Gly Val Tyr Thr Lys Val  
450 455 460

Thr Asn Tyr Leu Gly Trp Ile Arg Asp Asn Met His Leu  
465 470 475